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- **②** Designated Contracting States AT BE CHIDE DK ES FRIGBIGRIE IT LI LU MC
- ③ Applicant ONO PHARMACEUTICAL CO., LTD. Kyoto-shi Sakyo-ku, Kitashirakawa Olwakecho Kan'yuchi, Applicant: Honjo, Tasuku Osaka (JP) Osaka-shi Chuo-ku 1-5, Doshomachi 2-chome
- (2) Inventor: Honjo, Tasuku 19, Eikando Nishimachi 102, Fujii Eikando Bunko, Inventor: Shinohara, Takashi Massachusotts 02164 (JP) Sakyo-ku Kitashirakawa Olwakecho, Newton, 35 Cottage Street Inventor: ishida, Yasumasa Kyoto (JP) Kyoto-shi, Kan'Yuchi,
- 3 Representative: Henkel, Feller, Hänzel & D-81675 München (DE) Möhistrasso 37 Partner

Kyoto-shi Sakyo-ku,

Kypto (JP)

(a) A novel poptide related to human programmed cell death and DNA enceding it.

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(e) A membrane protein related to human programmed cell death (PD-1) and DNA encoding the said protein is new PD-1 protein may be useful for the treatment of various infections, immunological depression or acceleration, or tumour etc.

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Field of the inventior

The present invention is related to a novel peptide cell death and DNAs encoding it.

no have been shown to require de novo synthesis of RNA and protein. accidental deaths that occur by pathological mechanisms. Most of the cells undergoing programmed death various species of animals. Such cell deaths are generally considered 'programmed' and distinguished from Developmentally and physiologically controlled cell deaths can be observed in almost all tissues of

programmed cell death. These facts suggest that at least a few genes, if not specified ones, must be expressed to cause

while the mitochondria and other organelles are unaffected. A unique biochemical feature of apoptotic cells of cell death. In cells dying by apoptosis, the chromatin condenses around the periphery of the nucleus includes fragmentation of DNA into oligonucleosomal pieces. In mammats, apoptosis is often associated with programmed cell death morphologically and biochemically, but some of the cells undergoing proapoptotic cell deaths that can be induced in the absence of any protein synthesis. grammed death apparently do not show the characteristic teatures of apoptosis. In addition, there are The term 'apoptosis', on the other hand, is used to describe the morphological characteristics of a class

Thus, it is important to note that apoptosis is not synonymous with programmed cell death. Recently, if has been apparent that bct-2 which is a oncogene and in mortalized B cells by protection

the cell death, it was shown the importance to control the cell death

Related Arts

Kyoto (JP)

From now, certain peptides which are related to programmed cell death were reported. In such peptides, one of the representative is Fas antigen (tloh, N. et al., Cell, 68, 233 (1991)). Human Fas antigen is a polypeptide consisting from 335 amino acids, having signal peptide consisting

႘ divided to extracellular domain (157 amino acids), transmembrane region (17 amino acids) and cytoplasmic 16 hydrophobic amino acids N-terminal and it was considered that its mature protein have a structure inducing cell death. domain (145 amino acids). And it was thought that Fas anligen had a function of receptor to a factor (ligand)

Purpose of the invention

represented by Fas antigen. The purpose of the present invention is to find novel polypaptide which is alternative from polypaptides

have succeeded to find a quite novel polypeptide and DNA encoding it, and then completed the present sequence have been decided and its amino acid sequence have been deduced. And the present inventors in this invention, gene deeply related to programmed cell death have been isolated, its nucleotide

mouse T cell hybridoma 2B 4.11 (Japanese Patent Kokai 5-336973) was used as probe. To isolate a gene deeply related to programmed cell death in human, mouse PD-1 which obtained from

that of the polypeptide of the present invention except for mouse PD-1, when amino acid sequences of the that the polypeptide has no homology to Fas antigen. sequences in data base of National Biomedical Research Foundation, Needless to say, it was confirmed polypeptide identified in the present invention was compared by a computer program for all known There was no polypeptides having amino acid sequence which is identical to or has high homology to

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Constitution of the Invention

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(abbreviated human PD-1 hereafter). The present invention is related to polypeptide which is deeply related to programmed cell death

in substantially purified form, a homologue thereof or a tragment of the sequence or homologue of fragment, and DNA encoding such a polypeptide. More particularly, the present invention is related to DNA having the nucleotide sequence shown in SEQ. ID. No. 2 or 3, and DNA having a fragment which is selectively hybridizing to nucleotide sequence shown in SEQ. ID. No. 2 or 3. The present invention is concerned with a polypeptide having the amino acid shown in SEQ. ID. No. 1,



The present invention is related to:

- (1) a polypeptide having an amino acid sequence shown in SEQ. ID. No. 1,
 - (3) a DNA having a nucleotide sequence shown in SEQ. ID. No. 2, and (2) a DNA encoding the polypeptide described above (1).
 - (4) a DNA having a nucleotide sequence shown in SEQ. ID. No. 3.

A polypeptide of SEQ. ID. No. 1 in substantially purified form will generally comprise the polypeptide in a preparation in which more than 90%, e.g. 95%, 98% or 99% of the polypeptide in the preparation is that of the SEQ. ID. No. 1.

80% and more preferably at least 95% homologous to the polypeptide of SEO, ID. No. 1 over a region of at least 20, preferably at least 30, for instance 40, 60 or 100 more contiguous amino acids. Such polypeptide A polypeptide homologue of the SEQ. ID. No. 1 will be generally at least 70%, preferably at least 80 or homologues will be referred to below as a polypeptide according to the invention.

example 20, 25, 30, 40, 50 or 60 amino acids in length, and are also encompassed by the term "a Generally, fragments of SEQ. ID. No. 1 or its homologues will be at least 10, preferably at least 15, for polypeptide according to the invention" as used herein.

A DNA capable of selectively hybridizing to the DNA of SEQ. ID. No. 2 or 3 will be generally at least 70%, preferably at least 85 or 90% and more preferably at least 85% homologous to the DNA of SEQ. ID. No. 2 or 3 over a region of at least 20, preferably at least 30, for instance 40, 60 or 100 or more contiguous nucleotides. Such DNA will be encompassed by the term "DNA according to the invention".

Fragments of the DNA of SEQ. ID. No. 2 or 3 will be at least 15, preferably at least 20, for example 25, 30 or 40 nucleatides in length, and are also encompassed by the term "DNA according to the invention" as used herein. 8

of the promoter. The vector may contain one or more selectable marker genes, for example a ampicillin resistance gene. The vector may be used in vitro, for example of the production of RNA corresponding to A further embodiment of the invention provides replication and expression vectors comprising DNA according to the invention. The vectors may be, for example, plasmid, virus or phage vectors provided with an origin of replication, optionally a promoter for the expression of the said DNA and optionally a regulator the DNA, or used to transfect or transform a host cell. 2

A further embodiment of the invention provides host cells transformed or transfected with the vectors for the replication and expression of DNA according to the invention, including the DNA SEQ. ID. No. 2 or 3 or the open reading frame thereof. The cells will be chosen to be compatible with the vector and may for example be bacterial, yeast, insect or mammalian. 8

A further embodiment of the invention provides a method of producing a polypeptide which comprises culluing host cells of the present invention under conditions effective to express a polypeptide of the invention. Preferably, in additions such a method is carried out under conditions in which the polypeptide of 23

the invention is expressed and then produced from the host cells.

DNA according to the invention may also be inserted into the vectors described above in an antisense orientation in order to proved for the production of antisense RNA. Antisense RNA may also be produced by synthetic means. Such antisense RNA may be used in a method of controlling the levels of a polypeptide of the invention in a celi. \$

The invention also provides monoclonal or polyclonal antibodies to a polypaptide according to the Invention. The Invention further provides a process for the production of monocional or polyclonal antibodies to the polypeptides of the Invention. Monocional antibodies may be prepared by conventional hybridoma. technology using a polypeptide of the invention or a fragment thereof, as an immunogen. Polycional antibodies may also be prepared by conventional means which comprise inoculating a host animal, for oxample a rat or a rabbit, with a polypeptide of the invention and recovering immune serum. ş

The present invention also provides pharmaceutical compositions containing a polypeptide of the invention, or an antibody thereof, in association with a pharmaceutically acceptable diluent and/or carrier,

The polypeptide of the present invention includes that which a part of their amino acid sequence is lacking (e.g., a polypeptide comprised of the only essential sequence for reveating a biological activity in an amino acid sequence shown in SEO. ID. No.1), that which a part of their amino acid sequence is replaced by other amino acids (e.g., those replaced by an amino acid having a similar property) and that which other amino acids are added or inserted into a part of their amino acid sequence, as well as those having the amino acid sequence shown in SEQ. ID. No. 1. 8

As known well, there are one to six kinds of codon as that encoding one amino acid (for example, one kind of codon for Methioine (Met), and six kinds of codon for feucine (Leu) are known). Accordingly, the nuclootido sequence of DNA can be changod in order to encode the polypeptide having the same amino 8

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The DNA of the present invention, specified in (2) includes a group of every nucleotide sequences encoding polypeptides (1) shown in SEQ. ID. No. 1 . There is a probability of improving a yield of production of a polypeptide by changing a nucleotide sequence.

The DNA specified in (3) is the embodiment of DNA shown in (2), and is sequence in the natural form.

The DNA shown in (4) indicates the sequence of the DNA specified in (3) with a non-translational

The DNA of the present Invention may be obtained by gene recombination, chemical synthesis known methods for the skilled in the arts.

Human PD-1 includes a series of polypeptides which are deferent from Fas antigen in structural feature and commonly in mammals. That is, PD-1 of the present invention includes human PD-1 declared in the present invention and PD-1 of the other mammals which have high homology (it means immunological equivalent which can be cross-reacted to human PD-1 antigen).

The structural feature of human PD-1 is as follows:

Human PD-1 is predicted a membrane binding type protein consisting with 288 amino acids, it contains two hydrophobic regions, one at the N terminus and the other in the middle, which are likely to serve as a signal peptide and a transmembrane segment, respectively. 5

Comparison of the N-terminal sequence of the PD-1 protein with typical signal peptide cleavage sites suggests that the signal peptide would be from Met1 to Arg20. Thus, the predicted mature form of the PD-1 protein would contain 268 amino acids and consists of an extracellular domain (147 amino acids), a transmembrane region (27 amino acids) and a cytoplasmic domain (94 amino acids). Four potential Nglycosylation sites are found in the putative extracellular domain. 2

Comparison of the amino acid sequence of the PD-1 protein with all sequences registered in the National Biomedical Research Foundation data base revealed that the extracellular domain of the PD-1 protein is homologous to some members of the immunoglobulin superfamily, immunoglobilin domains havo been classified into V, C1 and C2 sets based on the conserved amino acid patterns and the number of

antiparallel beta-strands. The 68 amino acid residue between two cystein residues (Cys54 and Cys123) in PD-1 bear resemblance to a disulfidelinked immunogloblin domain of the V-set sequences. In addition, all of the four amino acid residues characteristic of many V-set sequences are also conserved in PD-1 (Arg94, Phe95, Asp117 and Gly119). 5

The cytoplasmic domain of the predicted PD-1 protein contains a variant form of the consensus sequence (Asp/Glu-X8-Asp/Glu-X2-Tyr-X2-Leu/ile-X7-Tyr-X2-Leu/ile) found in the cytoplasmic talls of most of the polypeptides associated with antigen receptors and Fc receptors. It was recently shown that one signal unit of this consensus sequence is sufficient to transduce signals. 8

It is thought that PD-1 of other mammats would be similar to human PD-1 in structural feature, whethr or not, number or kinds of its amino acid would be different in its sequence. ĸ

Invention may be obtained by chemical synthesis, by PCR method or by hybridization making use of a fragment of DNA of the present invention, as probe. Furthermore, DNA of the present invention may be DNA encoding human PD-1 of the present invention may be prepared by the following method. Once the nucleotide sequences shown in SEO. ID. Nos. 2 and 3 are determined, DNA of the present obtained in a desired amount by transforming with a vactor DNA inserted a DNA of the present invention into a proper host, followed by culturing the transformant. \$

The PD-1 polypeptides of the present invention (shown in SEQ. ID. No. 1) may be prepared by:

(1) isolating and purifying from an organism or a cultured cell, (2) chemically synthesizing, or

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using a skill of biotechnology.

preferably, by the method described in (3).

Examples of expression system when preparing a polypeptide by using a skill of biotechnology, is, for example, the expression system of bacteria, yeast, insect cell and mammalian cell. 20

of a DNA encoding mature peptide, connecting the DNA thus obtained to the downstream of a proper promoter (e.g., trp promoter, lac promoter, NPL promoter, 77 promoter etc.), and then inserting it into a vector (e.g., pBR322, pUC18, pUC19 etc.) which functions in an E. coli strain to prepare an expression For example, the expression in E. coli may be carried out by adding the initiation codon (ATG) to 5' end 99

Then, an E. coll strain (e.g., E. coll DH1 strain, E. coll JM109 strain, E. coll HB101 strain, etc.) which is transformed with the expression vector thus obtained may be cultured in a proper medium to obtain the

desired polypeptide. When a signal peptide of bacteria (e.g., signal peptide of pel B) is utilized, the desired polypeptide may be also released in periplasm. Furthermore, a fusion protein with other polypeptide may be also produced easily.

- DNA encoding PD-1 into the downstream of a proper promoter (e.g., SV40 promoter, LTR promoter Furthermore, the expression in a mammalian cell may be carried out, for example, by inserting the total
- medium. The polypeptide thus obtained may be isolated and purified by conventional blochemical methods of other animals as proba. obtained, and then culturing the transformant in a proper medium to get a desired polypeptide in the culture (e.g., mankey COS-7 cell, Chinase hamster CHO cell, mouse L cell etc.) with the expression vector thus virus vector, SV40 vector, etc.) to obtain an expression vector, and transforming a proper mammalian cell metallothionein promoter etc.) in a proper vector (e.g., retrovirus vector, papilloma virus vector, veccinis DNA encoding PD-1 gene obtained by the present invention may be used for the isolation of PD-1 gene
- The cDNA having a nucleotide sequence shown in SEO. ID. No. 3 may be prepared according to the
- (ii) by preparing first strand (single stranded DNA) from mRNA thus obtained, followed by preparing human esophageal cancer cell line), (i) by isolating mRNA from a cell line which products the polypeptide of the present invention (e.g.
- second strand (double stranded DNA) (synthesis of cDNA).
- (iii) by Inserling cDNA thus obtained into a proper phage vector, (v) by transforming host cells with the recombinent DNA thus obtained (preparation of cDNA library). (v) by screening with plaque hybridization from cDNA library thus obtained with cDNA of mouse PD-1 as
- (vi) by preparing phage DNA from positive clone obtained, subcloning cDNA released into plasmid
- complete length by combining them. vector, proparing restriction enzyme map, and (vii) by deciding sequence of each restriction enzyme fragment, and by obtaining the full sequence of

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- of the present invention is preferably human cell line YTC3. (described in Methods in Enzymology, vol. 154, p 3, 1987) from human cell line after stimulation by a proper stimulant (e.g., IL-1 etc.) or without stimulation. Examples of the cells which product the potypeptide Explained in detail, step (i) may be carried out in accordance with the method of Okayama, H. et al.
- may be preferably used. As examples of the plasmid vector used in the step (iii), many plasmid vectors (e.g., pBR 322, pBluescript II) and phage vectors (e.g., \(\chi_2\)), \(\chi_3\) is the known, and phage vector \(\chi_3\) (43.3 kbp.Stratagene) accordance with the method of Gubler & Hoffman (Gene, vol. 25, pp. 263, 1983) with a slight modification Steps (ii), (iii) and (iv) are a series of steps for preparing cDNA library, and may be carried out in
- As host cell used in step (iv), E. Coll NM514 (Strategene) may be preferably used.
- The stops (v) and (vi) may be carried out in accordance with the method described in Molecular Cloning (Sambrook, J., Fritsh, E. F., and Maniatis, T. Cold Spring Harbor Laboratory Press (1989)).
- The step (vii) may be carried out in accordance with the method described in Molecular Cloning (written by Sambrook, J., Fritsch, E. F. and Maniatis, T., published by Cold Spring Harbor Laboratory Press in
- the dideaxy termination method. The sequencing in the step (vii) may be carried out in accordance with the method of Maxam-Gilbert or
- length. The confirmation may be carried out by Northern analysis with the said cDNA as probe (see almost same longth of mRNA obtained in the hybridizing band. Molecular Cloning described bolore). It is thought that cDNA is almost complete length, if length of cDNA is It is necessary to examine whether or not the cDNA thus obtained codes complete or almost complete
- or for the purpose of diagnosing diseases, and the like. polypoptide and protection mechanism in living organism, immunological function or diseased like tumour primer and thereby, and may be utilized for the purpose of investigating the relationship between the said DNA or DNA fragments encoding PD-1 gene may be used for detection of PD-1 gene as probe or a
- thereof which are expected to possess various use. by conventional gone recombination the PD-1 polypeptide, polypeptide fragment thereof or derivatives The DNA of the present invention may be utilized as an important and essential template in preparing
- It is expected that the polypeptide, fragment polypeptides thereof or derived polypeptides thereof may
- be used for the treatment of infections, depression or acceleration of immunological function or tumour.

 Further, polycional and monocional antibody against the polypoptide or polypoptide fragments of the
 prosent invention can be prepared by conventional method, and they can be used to quantilate the said

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polypeptide in organism, and thereby, may be utilized for the purpose of investigating the relationship between the said polypeptide and diseases, or for the purpose of diagnosing diseases, and the like. The said monocional antibody per se, chimeric antibody against human antibody may be used for the treating

Polyclonal and monoclonal antibody thereof may be prepared by conventional methods by using the said polypeptide or the fragment thereof as an antigen

- intravenously or intraventricularly. administered systemically or partially, usually by oral or parenteral administration, preferably orally For the purpose of the present invention, the polypeptide of the present invention may be normally
- doses per person per dose are generally between 100 ug and 100 mg, by oral administration, up to several The doses to be administered are determined depending upon age, body weight, symptom, the desired therapeutic effect, the route of administration, and the duration of the treatment etc. In the human adult, the times per day, and between 10 ug and 100 mg, by parenteral administration up to several times per day.
- in which doses lower than or greater than the ranges specified above may be used.

 Administration of the compounds of the present Invention, may be as solid compositions, liquid As mentioned above, the doses to be used depend upon various conditions. Therefore, there are cases
- compositions or other compositions for oral administration, as injections, liniments or suppositories etc.
- powders, granules. Capsules include soft capsules and hard capsules.

 In such compositions, one or more of the active compound(s) is or are admixed with at least one inert Solid compositions for oral administration include compressed tablets, pills, capsules, dispersible
- dituent (such as tactose, mannitol, glucose, hydroxypropyl cellulose, microcrystalline cellulose, starch, stearate etc.), disintegrating agents (such as cellulose calcium glycolate, etc.), stabilizing agents (such as human serum albumin, lactose etc.), and assisting agents for dissolving (such as arginine, asparaginic acid normal practice, additional substances other than inert diluents: e.g. lubricating agents (such as magnesium polyvinylpyrrolidone, magnesium metasilicate aluminate, etc.). The compositions may also comprise, as is
- than two films. And further, coating may include containment within capsules of absorbable materials such getatin, hydroxypropyl cellulose or hydroxypropylmethyl cellulose phthalate, etc.), or be coated with The tablets or pills may, if desired, be coated with a film of gastric or enteric material (such as sugar more
- ដ flavouring agents, perfurning agents, and preserving agents.

 Other compositions for oral administration included spray compositions which may be prepared may also comprise adjuvants (such as wetting agents, suspending agents, etc.), sweetening agents diluent(s) commonly used in the art (purified water, ethanol etc.). Besides inert diluents, such compositions syrups and elixirs. In such compositions, one or more of the active compound(s) is or are contained in inert Liquid compositions for oral administration include pharmaceutically-acceptable emulsions, solutions,
- in their entireties by reference) may be used. buffer (sodium chloride, sodium citrate, citric acid, etc.). For preparation of such spray compositions, for example, the method described in the United States Patent No. 2,868,691 or 3,095,355 (herein incorporated comprise additional substances other than inert diluents: e.g. stabilizing agents (sodium sulfite etc.), isotonic known methods and which comprise one or more of the active compound(s). Spray compositions may
- diluents(s)(propylene glycpl, polyethylene glycol, plive oil, ethanol, POLYSOLBATE 80 TM, etc.) inent aqueous diluent(s) (distilled water for injection, physiological salt solution, etc.) or inent non-aqueous and emulsions. In such compositions, one or more active compound(s) is or are admixed with at least one Injections for parenteral administration include sterile aqueous or non-aqueous solutions, suspensions
- assisting agents such as assisting agents for dissolving (arginine, asparaginic acid, etc.). hjections may comprise additional other than inert dituents: e.g. preserving agents, wetting agents, emulsifying agents, dispersing agents, stabilizing agent (such as human serum abumin, tactose, etc.), and
- solid compositions, for example, by freeze-drying, and which can be dissolved in sterile water or some other sterile diluents for injection immediately before use. sterilizing agents in the compositions or by irradiation. They may also be manufactured in the form of sterile They may be sterilized for example, by filtration through a bacteria-retaining filter, by incorporation of
- (ointment, etc.), suppositories for rectal administration and pessaries which comprise one or more of the Other compositions for parenteral administration include liquids for external use, and endermic liniments

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active compound(s) and may be prepared by known methods.

Examples

The following examples are illustrated, but not limit, the present invention.

Example 1: Cell culture

The human cell lines (CESS, HPB-ALL, Jarkat, TC3, CCRF-CEM, JM and MOLT-4F) were cultured in RPMI 1840 (Gibco) supplemented with 10% heat-inactivated fetal call sorum, 2mM glutamine, 50 µM 2mercaptoethanol, 100U/ml penicillin and 100 ug/ml streptnycin 2

Example 2: Northern blot analysis

at 80 °C for 2 hrs. Random priming was carried out to the Eco-RI fragment (1kb) containing the coding region of mouse PD-1, with ²²⁹ to prepare as probe. Specific activity of this probe was about 9 x 10* Total RNA was prepared from indicated cell lines by extraction with guanidium isothlocyanate method (see molecular cloning described before), and poly (A)* RNA was isolated from the total RNA by oligotex dT 30 (super) (Daiichi Chemical Co.), 3 ug of poly (A)* RNA was separated on a 1,2% formaldehydeagarose gel, and transferred to a nylon membrane (Blodyne A, Japan Genetic). The filter was baked d.p.m./ug. Hybridization was carried out in 10 x Denhardi's, 1M NaCl, 50mM Tris (pH 7.5), 10 mM EDTA 1% SDS and 1 mg/ml soniceled salmon sperm DNA at 65 °C for 15 hrs. The filter was washed in 1 x SSC 0.1% SDS at 65 °C for 10 mins. Hybridization signal (2.3 kb) was observed from lymphocyte cell line YTC3 5 2

Example 3: Construction of cDNA library and cloning of human PD-1 cDNA

SK plasmid vector (Stratagene) was used as probe. The filter was washed with 1 x SSC and 0.1% SDS at 60 · C for 10 mins. 51-Positive signals were observed from 1.2 x 10⁶ phages by autoradiography. These clones were purified to single. Further analysis was carried out about 23 clones picked up, the longest Saver cDNA Synthosis Kit (Pharmacia), Synthesis of litst strand cDNA was carried out with oligo of primer. Double stranded cDNA which was ligated EcoRI-Notl adopter, was cloned into \(\text{g} \) 10 vector, and packaged into phage (Gigapack II Gold, Stratagene). Phage was plated on a lawn of E. Coll NMS14, Phage DNA was cDNA library was constructed with 5 ug of poly (A)* RNA extracted from YTC3 cell lines by using Time transfected to duplicated library filters from each plate. The filters were baked at 80 °C for 2 hrs and hybridized at 60 °C for 15 hrs. The mouse PD-1 coding region (1 kb) excised with EcoRI from Bluescript cDNA insert observed was 2.1 kb. This result coincided to the result of Southern Blot analysis. 25

Example 4: Sequencing of DNA

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Amersham). Specific primer of Bluescript plasmid was used as a sequencing primer. Nucleotide sequencing was actured out by fully sequencing for both strands of the CDA. From the result, incleotide sequence shown in Sec. 10. L. 2 was obtained. Deduced peptide sequence (shown in SEC. ID. No. 1) was determined from the nucleotide sequence. Total number of deduced amino acid is 288, and the number is the same as The cDNA inserts isolated from human cDNA library were subcloned into Bluescript SK plasmid vectors (Stratagene), and sequenced by the dideoxynuclectide chain termination method (Sanger et al., 1977) using a modified T7 DNA polymerase (United States Biochemical) and [a-39jdCTP (3000 Ci/mmol that of mouse PD-1. Homology of was found about 60 % each other. \$

Example 5 : Southern Blotting 8

described before). DNAs were digasted by EcoRI, BamHI or hind III followed according to the manufactures-recommendation, isolated by electrophorasis (100V, 0.8% agarose gel, TEA buffer). DNA fragments were washed with 0.25N HCl for 10 mins, denatured with 0.2N NaOH / 0.6M NaCl for 30 mins, neutralized with 0.8M NaCl / 0.2M Tris (pH 7.5) for 1 hz., transferred to nylon membrane (Bidyne A) which is used Genomic DNA were isolated from kinds of animal cells by conventional method (see Molecular Cloning standard Southern procedure. The filter were baked for 2 hrs. ş

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Random priming was carried out to the EcoRI-Stul fragment (900 bp) containing the coding region of Hybridization was carried out in 10 x Denhardt's, 1M NaCl, 50mM Tris (pH 7.5), 10 mM EDTA, 1% SDS and at 65 °C for 10 mins. Only one band was detected by autoradiography when the clone was cut with any enzyme, it was found that human PD-1 gene exists as single copy. human PD-1, 32P to prepare as probe. Specific activity of this probe was about 9 x 10⁸ d.p.m./lug. Img/ml sonicated salmon sperm DNA at 65 °C for 10 mins. The filter was washed in 1 x SSC, 0.1% SDS

(hybridization and washing) described in example 2, using EcoRI fragment (1kb) containing coding region of mouse PD-1 as probe. Hybridization signals were detected from only genomic DNA of mouse and human, and were not detected from genomic DNA of Drosophila, Xenopus and rabbit. Southern hybridization was carried out with genomic DNA of kinds of animals by the same condition

Example 6: Isolation of genomic clone of human PD-1

A genomic DNA library from esophageal cancer cell line was constructed in the x DASH il vector via Sau3Al partial digestion and ligation into the BamH site (obtained from Dr. Nishiyama, 1st Dapt. of Pathology, School of Medicine, Kyoto University). The human PD-1 gene was isolated from this fibrary by hybridization with the human PD-1 total cDNA excised with EcoRI digestion from Bluescript SK vector. The probe was labeled with ^{32P} by random priming. Two positive clones was Isolated and puritied from 1 x 10^t phage plaque, digested by several restriction enzymes, and analyzed by Southern hybridization using the same probe. From CISS (chromosomal in atlu suppression) , it was found that human PD-1 gone was

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SEQUENCE LISTING

(1) GENERAL INFORMATION:

(2) INFORMATION FOR SEQ ID NO:1: (1v) COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk (B) COMPUTER: BLM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS (D) SOFTMARE: Patantin Release #1.0, Version #1.30 (EPO) (111) NUMBER OF SEQUENCES: 4 (11) TITLE OF INVENTION: A NOVEL PEPTIDE RELATED TO HUMAN PROGRAMMED CELL DEATH AND DNA ENCODING IT (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: (11) MOLECULE TYPE: protoin (1) SEQUENCE CHARACTERISTICS: (A) LENGTH: 288 maino acids (B) TYPE: maino ncid (C) STRANDEDNESS: single (D) TOPOLOGY: linear Asn Pro Pro Thr Pho Ser Pro Ala Leu Leu Val Val Thr Glu Gly Asp 35 40 45 Lou Gly Trp Arg Pro Gly Trp Phe Leu Asp Ser Pro Asp Arg Pro Trp 20 25 Met Gln Ile Pro Gln Ala Pro Trp Pro Val Val Trp Ala Val Leu Gln 15 (1) APPLICANT: (A) NAME: ONO PHARHACEUTICAL CO., LTD. (B) STREET: 1-5, Doshomschi 2-chome (C) CITY: Chuo-ku, Osaka-shi (D) STATE: Osaka (E) COUNTRY: Japan (F) POSTAL CODE (ZIP): 541 (A) NAME: HONJO, TASUKU (B) STREET: Kan'yuchi, Kitashirakawa Olwakecho, Sakyo-ku (C) CITY: Kyoto-shi . (D) STATE: Kyoto (E) COUNTRY: Japan (F) POSTAL CODE (EIP): 606

Net Gly Thr Ser Ser Pro Ala Arg Arg Gly Ser Ala Asp Gly Pro Arg 260 260 270 Ser Ala Gln Pro Leu Arg Pro Glu Asp Gly His Cys Ser Trp Pro Leu 275 280 Cys Val Pro Glu Gln Thr Glu Tyr Ala Thr Ile Val Phe Pro Ser Gly 245 Glu Leu Asp Phe Gln Trp Arg Glu Lys Thr Pro Glu Pro Pro Val Pro 225 230 230 Leu Lys Glu Asp Pro Ser Ala Val Pro Val Pha Ser Val Asp Tyr Gly
210 225 220 Ser Arg Ala Ala Arg Gly Thr Ile Gly Ala Arg Arg Thr Gly Gln Pro Arg Ser Ala' Gly Gln Phe Gln Thr Leu Val Val Gly Val Val Gly Gly 175 175Thr Glu Arg Arg Ala Glu Val Pro Thr Ala His Pro Ser Pro Ser Pro 145 150 150 160 Ala Pro Lys Ala Gln Ile Lys Glu Ser Leu Arg Ala Glu Leu Arg Val 130 Ala Arg Arg Asn Asp Ser Gly Thr Tyr Leu Cys Gly Ala Ile Ser Leu 115 Val Thr Gln Leu Pro Asn Gly Arg Asp Phe His Met Ser Val Val Arg 100 105 Lou Leu Gly Sor Leu Val Leu Leu Val Trp Val Leu Ala Val Ile Cys
180
180 Ala Phe Pro Glu Asp Arg Ser Gln Pro Gly Gln Asp Cys Arg Phe Arg 95 Leu Asn Trp Tyr Arg Met Ser Pro Ser Asn Gln Thr Asp Lys Leu Ala $65 \ \ \, 70 \ \ \, 75 \ \ \,$ Asn Ala Thr Phe Thr Cys Ser Phe Ser Asn Thr Ser Glu Ser Phe Val $50\,$

(2) INFORMATION FOR SEQ ID NO:2:

(1) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 864 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(11) MOLECULE TYPE: CDNA to mRNA

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(x1) SEQUENCE DESCRIPTION, SEQ ID NO:2:		244004
ATGCAGATCC CACAGGCGCC CTGGCCAGTC GTCTGGGCGG TGCTACAACT GGGCTGGCGG	09	
CCAGGATGGT TETTAGACTE CECAGACAGG CECTGGAACE CECECAGETT CTECECAGE	120	
CTGCTCGTGG TGACCGAAGG GGACAACGCC ACCTTCACCT GCAGCTTCTC CAACACATCG	180	
GAGAGCTICG TGCIAAACIG GTACCGCATG AGCCCCAGGA ACCAGAGGA CAAGCTGGCC	240	CAGATO
GCCTTCCCCG AGGACCGCAG CCAGCCCGGC CAGGACTGCC GCTTCCGTGT CACACAACTG	300	
CCLANCGGGC GTGACTTTCCA CATGAGCGTG GTCAGGGCCC GGCGCAATGA CAGCGGCACC	360	
TACTICTGTG GGGCCATCTC CCTGGCCCCC AAGGCGCAGA TCAAAGAGAG CCTGCGGGCA	420	TCCCGG
GAGTICAGGG TGALAGAGAG AAGGGCAGAA GTGCCCACAG CCCACCCCAG CCCTCACCC	480	CCCTCAC
AGGTCAGCCG GCCAGTTCCA AACCCTGGTG GTTGGTGTCG TGGGCGGCCT GCTGGGCAGC	540 15	
CTGGTGCTGC INGTCTGGGG CCTGGGGGTC ATCTGCTCCC GGGCGGCACG AGGALAAIA	009	TTTCCT
GGAGCCAGGC GCACCGGCCA GCCCTGAAG GAGGACCCCT CAGCCGTGCC TGTGTTCTCT	999	AGTGCCC
GIGGACTATG GGGAGCIGGA TITUCAGIGG CGAGAGAAGA CCCCGGAGCC CCCCGIGCCC	720 20	•
TGTGTCCCTG AGCAGAGGGA GTATGCCACC ATTGTCTTTC CTAGCGGAAT GGGCACCTCA	780	(2) INE
TCCCCCCCC GCAGGGCTC AGCTGACGGC CCTCGGAGTG CCCAGCCACT GAGGCCTGAG	840	Ξ
GATGGACACT GCTCTTGGCC CCTC	864 25	
(2) INFORMATION FOR SEQ ID NO:3:		
(1) SEQUENCE CHARACTERISTICS:		77)
(A) LENGTH: 911 base pairs (B) TYPE: nucleic cald (C) STRANDEDNESS: single (D) TOPOLOGY: lines	8	
(11) MOLECULE TYPE: CDNA to mRNA		.
	50	
(x1) SEQUENCE DESCRIPTION: SEQ ID NO.3:		(
CACTETGGTG GGGTGCTCC AGGCATGCAG ATCCCACAGG CGCCCTGGCC AGTCGTCTGG	09	
GCGGTGCTAC AACTGGGCTG GCGGCCAGGA TGGTTCTTAG ACTCCCCAGA CAGGCCTGG	120	(1
AACCCCCCCA CCTTCTCCCC AGCCCTGCTC GTGGTGACCG AAGGGGACAA CGCCACCTTC	180	
ACCIGEAGCI ICICCAACAC AIGGAGAGC TICGIGCIAA ACIGGIACCG CAIGÁGCCC	240	
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	TGCGGCTTCC GTGTCACACA ACTGCCCAAC GGGCGTGACT TCCACATGAG CGTGGTCAGG 360
40	GCCGGGGGA ATGACAGGGG CACCTACCTC TGTGGGGCCA TCTCCCTGGC CCCCAAGGGG
	CAGATCANAG AGAGCCTGCG GGCAGAGCTC AGGGTGACAG AGAGNAGGGC AGAAGTGCCC 480
	ACAGCCCACC CCAGCCCCTC ACCCAGGTCA GCGGCCAGT TCCAAACCCT GGTGGTTGGT 540
5	STCGTGGGCG GCCTGCTGGG CAGCTAGGTCT GGGTCCTGGC CGTCATCTGC 600
	TCCCGGGCCG CACGAGGGA AATAGGAGCC AGGCGCACCG GCCAGCCCCT GAAGGAGGAC 660
	CCCTCAGCCG TGCCTGTGTT CTCTGTGGA. TATSGGGAGC TGGATTTCCA GTGGCGAGAG 720
15	AAGACCCCGG AGCCCCCGT GCCCTGTGTC CCTGAGCAGA CGGAGTATGC CACCATTGTC 780
	TITCCTAGGG GAATGGGCAC CICAICCCC GCCCGCAGGG GCTCAGCIGA GGCCCTCGG 840
	AGTGCCCAGG CACTGAGGCC TGAGGATGGA CACTGCTCTT GGCCCCTCTG ACCGGCTTCC 900
8	TTGGCCACCA GAGTTCTGCA G
	(2) INFORMATION FOR SEQ ID NO:4:
25	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 921 base pairs (B) TYPE: nucleic acid (C) STRANDENESS: single (D) TOPOLOGY: linear
	(11) MOLECULE TYPE: CDNA to mRNA
8	(vi) ORIGINAL SOURCE: (A) ORGANISM: HOme sapiens (H) CELL LINE: YTC3
ĸ	(ix) FEATURE: (A) NAME/FEY: CDS (B) LOCATION: 25.888 (C) IDENTIFICATION METHOD: P
\$	(4x) FEATURE: (A) RAMENTON: 25.84 (B) LOCATION: 25.84 (C) IDENTIFICATION METHOD: S
	(4x) PEATURE: (A) NAMENTER ED PEPTIDE (B) LOCATION: 05. 886 (C) IDENTIFICATION METHOD: S
ę	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

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627	GGA Gly	ATA 110 180	JP CV	615 999	yrg V93	33	GCC Ala 175	Pr 9	res DOL	cys Tec	I A	170 170	¥7 200	5.25	14A 315	in de	
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531	ren CJQ	Thr	GT ^W	Phe	145 679 679	61A 295	A1¤	TCA	95 V	94 95 95 95	Ser	200	Ser ≥	Pro	135 E E E	A14 CCC	ដ
48 3	The ACA	22.0 20.0	VAL OTO	100	25	35	AGA AGA	ត ខ្លួ	125 AC	14A	200	553	550	A14 120	λς 20 ξ	5.5	
435	AGC Ser	010 020	AAA Lys	PII SILV	55	71 939	Lys	944 233	¥1¥ 339	PET CING	10C	11e	105 200	520	Cyn	53	9
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	g	GG.	C1y GGC	Thr 230	Thr	VAI
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~ -		II 8		CCT	333	Asp.
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		TEACCGGCTT CCTTGGCCAC CAGTGTTCTG	Pro	A14	943	YE S
		TOTA	260 260	20,7	TAT	676
		63	GAT TAS	AGG Arg 245	919	Lys
	921	916	867	618	771	723

Claims

- A polypeptide having the amino acid sequence shown in SEQ. ID. No. 1 in substantially purified form, a homologue thereof or a fragment of the sequence or homologue of a fragment.
- 2. A polypeptide according to claim 1 having the amino acid sequence shown in SEQ. ID. No. 1.
- 30 3. DNA encoding a polypeptide according to claim 1.
- DNA according to claim 3 having the nucleotide sequence shown in SEQ. ID. No. 2 or a fragment thereot capable of selectively hybridizing to SEQ. ID. No. 2.
- DNA according to claim 3 having the nucleotide sequence shown in SEQ. ID. No. 3 or a fragment thereof capable of selectively hybridizing to SEQ. ID. No. 3.
- A replication and expression vector comprising DNA according to any one of claims 3 to 5.
- 7. Host cells transformed or transected with a replication and expression vector according to claim 6.
- A method of producing a polypeptide which comprises culturing host cells according to claim 7 under conditions effective to express a polypeptide according to claim 1 or 2.
- A monoclonal or polyclonal antibody to a polypeptide according to claim 1 or 2.

GCC AGG CGC ACC CGC CAG CCC CTC AAG GAC GAC CCC TCA GCC GTC CCT Ala Acg Acg The Gly Gln Pro Leu Lys Glu Asp Pro Ser Ala Val Pro 185

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A pharmaceutical composition containing a polypoptide according to claims 1 or 2 or an antibody according to claim 9 in association with a pharmaceutically acceptable dituent and/or carrier.

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Europálsches Patentamt European Patent Office

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£ Office européen des brevets

EUROPEAN PATENT APPLICATION

(12)

(88) Date of publication A3: 02.09.1998 Bulletin 1998/36

(51) Int. Ct.⁶: C12N 15/12, C07K 14/705, C12P 21/02, C07K 16/18, A61K 38/17, A61K 39/395

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(72) Inventors:

• Honjo, Tasuku
Kyoto-shi, Kyoto (JP)
• Ishida, Yasumasa
Newton, Massachusets 02164 (JP)
• Shinohara, Takashi
Sakyo-ku, Kyoto-shi, Kyoto (JP) (84) Designated Contracting States: AT BE CH DE DK ES FR GB GR IE IT LI LU MC NL PT SE (30) Priority: 01.03.1994 JP 55224/94

(71) Applicants:
ONO PHARMACEUTICAL CO., LTD.
Osaka-shi Osaka (JP)
Honjo, Tasuku
Sakyo-ku, Kyoto-shi, Kyoto (JP)

(74) Representative: Henkel, Feller, Hânzel Môhistrasse 37 81675 München (DE)

(54) A novel peptide related to human programmed cell death and DNA encoding it

(57) A membrane protein related to human programmed cell death (Pb.1) and DNA encoding the said protein is new. Pb.1 protein may be useful for the treatment of various intections, immunological depression or acceleration, or tumour etc.

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European Patent Office

EUROPEAN SEARCH REPORT

Application Number EP 95 10 2829

_	CLASSIFICATION OF THE APPLICATION (MI.C.I.E.)	OOOOAA			TECHNICAL FIELDS SEARCHED (IN.C.S)	C07K	7.	Ę	ourojian, u	Inhedon, or
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DOCHMENTS CONSIDE	Citation of document with undication, where appropriate, of relevant passages	ISHIDA Y ET AL: "Induced expression of the immunor gene superfamily, upon programmed death." EMBO 3. II (11) P3887-95 ENGLAND, XP002070368 • the whole document •	DATABASE NPI Section Ch, Week 9404 Derwent Publications Ltd. Class 804, AN 94-030912 & JP 05 336 973 A (HONJO Y) * abstract *	SHINOHARA I ET AL: "Structu chromosomal localization of gene (PDCD1)." GENOMICS. OCT 1994. 23 (3) P	STATES, XP000647607 * the whole document	i	The present search report has been drawn up for all claims	Puce of south	יוור וושפסר	CATEGORY OF CITED DOCUMENTS X particusty referant 4 team dates Y particusty referant 4 team dates Gocument of the same caregory Best processed to the same caregory O, sen-writter discount
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